Correction for Hraber et al., "Panels of HIV-1 Subtype C Env Reference Strains for Standardized Neutralization Assessments"

Peter Hraber,[a] Cecilia Rademeyer,[b] Carolyn Williamson,[b] Michael S. Seaman,[c] Raphael Gottardo,[d] Haili Tang,[e] Kelli Greene,[e] Hongmei Gao,[e] Celia LaBranche,[e] John R. Mascola,[f] Lynn Morris,[g] David C. Montefiori,[e] Bette Korber[a,h]

Theoretical Biology and Biophysics, Los Alamos National Laboratory, Los Alamos, New Mexico, USA[a]; Division of Medical Virology & Institute of Infectious Diseases and Molecular Medicine, University of Cape Town and NHLS, Cape Town, South Africa[b]; Center for Virology and Vaccine Research, Beth Israel Deaconess Medical Center, Harvard Medical School, Boston, Massachusetts, USA[c]; Vaccine and Infectious Disease Division, Fred Hutchinson Cancer Research Center, Seattle, Washington, USA[d]; Department of Surgery, Duke University Medical Center, Durham, North Carolina, USA[e]; Vaccine Research Center, National Institute of Allergy and Infectious Diseases, National Institutes of Health, Bethesda, Maryland, USA[f]; National Institute for Communicable Diseases, Johannesburg, South Africa[g]; New Mexico Consortium, Los Alamos, New Mexico, USA[h]

Volume 91, no. 19, e00991-17, 2017, https://doi.org/10.1128/JVI.00991-17. Supplemental material: Data Set S1 wrongly identifies the sequence MW_08_CH505_1_A2_KC247557 (CH0505.w4.3) as a transmitted-founder sequence. In row 184 ("KC247557"), column G ("is.TF"), "TRUE" should therefore be changed to "FALSE."

Revised supplemental material is posted at http://jvi.asm.org/content/91/19/e00991-17/suppl/DCSupplemental.